



## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/688,566

DATE: 06/17/2002

TIME: 19:34:49

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF3\06172002\I688566.raw

ENTERED

```

4 <110> APPLICANT: Lipovsek, Dasa
5      Wagner, Richard W
6      Kuimelis, Robert G
8 <120> TITLE OF INVENTION: PROTEIN SCAFFOLDS FOR ANTIBODY MIMICS
9      AND OTHER BINDING PROTEINS
12 <130> FILE REFERENCE: 50036/021004
14 <140> CURRENT APPLICATION NUMBER: US 09/688,566
15 <141> CURRENT FILING DATE: 2000-10-16
17 <150> PRIOR APPLICATION NUMBER: US 60/111,737
18 <151> PRIOR FILING DATE: 1998-12-10
20 <150> PRIOR APPLICATION NUMBER: US 09/456,693
21 <151> PRIOR FILING DATE: 1999-12-09
23 <150> PRIOR APPLICATION NUMBER: US 09/515,260
24 <151> PRIOR FILING DATE: 2000-02-29
26 <160> NUMBER OF SEQ ID NOS: 202
28 <170> SOFTWARE: FastSEQ for Windows Version 4.0
30 <210> SEQ ID NO: 1
31 <211> LENGTH: 122
32 <212> TYPE: DNA
33 <213> ORGANISM: Homo sapiens
35 <400> SEQUENCE: 1
36 ggaattccta atacgactca ctatagggac aattactatt tacaattaca atgcatcacc      60
37 atcaccatca cgtttctgat gttccgaggg acctggaagt tgttgctgcg acccccacca      120
38 gc                                                                    122
40 <210> SEQ ID NO: 2
41 <211> LENGTH: 104
42 <212> TYPE: DNA
43 <213> ORGANISM: Homo sapiens
45 <400> SEQUENCE: 2
46 ggaattccta atacgactca ctatagggac aattactatt tacaattaca atggtttctg      60
47 atgttccgag ggacctggaa gttgttgctg cgacccccac cagc                      104
49 <210> SEQ ID NO: 3
50 <211> LENGTH: 126
51 <212> TYPE: DNA
52 <213> ORGANISM: Homo sapiens
54 <220> FEATURE:
55 <221> NAME/KEY: misc_feature
56 <222> LOCATION: (1)...(126)
57 <223> OTHER INFORMATION: n = A,T,C or G
59 <221> NAME/KEY: misc_feature
60 <222> LOCATION: (1)...(126)
61 <223> OTHER INFORMATION: s = C or G
W--> 63 <221> misc_feature

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64 <222> LOCATION: 74, 75, 77, 78, 80, 81, 83, 84, 86, 87, 89, 90, 92, 93  
65 <223> OTHER INFORMATION: n = A,T,C or G  
W--> 67 <400> 3  
68 agcggatgcc ttgtcgtcgt cgtccttgta gtcgctcttc cctgtttctc cgtaagtgat 60  
W--> 69 cctgtaatat ctsnnsnnsn nsnnnsnnsn snnccagctg atcagtaggc tgggtgggggt 120  
70 cgcagc 126  
72 <210> SEQ ID NO: 4  
73 <211> LENGTH: 62  
74 <212> TYPE: DNA  
75 <213> ORGANISM: Homo sapiens  
77 <400> SEQUENCE: 4  
78 agcggatgcc ttgtcgtcgt cgtccttgta gtcgctcttc cctgtttctc cgtaagtgat 60  
79 cc 62  
81 <210> SEQ ID NO: 5  
82 <211> LENGTH: 99  
83 <212> TYPE: DNA  
84 <213> ORGANISM: Homo sapiens  
86 <400> SEQUENCE: 5  
87 ggaattccta atacgactca ctatagggac aattactatt tacaattaca atgcatcacc 60  
88 atcaccatca cctcttcaca ggaggaaata gccctgtcc 99  
90 <210> SEQ ID NO: 6  
91 <211> LENGTH: 132  
92 <212> TYPE: DNA  
93 <213> ORGANISM: Homo sapiens  
95 <220> FEATURE:  
96 <221> NAME/KEY: misc\_feature  
97 <222> LOCATION: (1)...(132)  
98 <223> OTHER INFORMATION: n = A,T,C or G  
100 <221> NAME/KEY: misc\_feature  
101 <222> LOCATION: (1)...(132)  
102 <223> OTHER INFORMATION: s = C or G  
W--> 104 <221> misc\_feature  
105 <222> LOCATION: 83, 84, 86, 87, 89, 90, 92, 93  
W--> 106 <223> OTHER INFORMATION: n = A,T,C or G  
W--> 108 <400> 6  
109 agcggatgcc ttgtcgtcgt cgtccttgta gtcgctcttc gtataatcaa ctccagggtt 60  
W--> 110 aaggccgctg atggtagctg tsnnnsnnsn snnaggcaca gtgaactcct ggacagggtt 120  
111 atttcctcct gt 132  
113 <210> SEQ ID NO: 7  
114 <211> LENGTH: 64  
115 <212> TYPE: DNA  
116 <213> ORGANISM: Homo sapiens  
118 <400> SEQUENCE: 7  
119 agcggatgcc ttgtcgtcgt cgtccttgta gtcgctcttc gtataatcaa ctccagggtt 60  
120 aagg 64  
122 <210> SEQ ID NO: 8  
123 <211> LENGTH: 101  
124 <212> TYPE: DNA  
125 <213> ORGANISM: Homo sapiens

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127 <400> SEQUENCE: 8
128 ggaattccta atacgactca ctatagggac aattactatt tacaattaca atgcatcacc      60
129 atcaccatca cctcttctat accatcactg tgtatgctgt c                          101
131 <210> SEQ ID NO: 9
132 <211> LENGTH: 114
133 <212> TYPE: DNA
134 <213> ORGANISM: Homo sapiens
136 <220> FEATURE:
137 <221> NAME/KEY: misc_feature
138 <222> LOCATION: (1)...(114)
139 <223> OTHER INFORMATION: n = A,T,C or G
141 <221> NAME/KEY: misc_feature
142 <222> LOCATION: (1)...(114)
143 <223> OTHER INFORMATION: s = C or G
W--> 145 <221> misc_feature
146 <222> LOCATION: 59, 60, 62, 63, 65, 66, 68, 69, 71, 72, 74, 75, 77, 78, 80,
147      81, 83, 84, 86, 87
148 <223> OTHER INFORMATION: n = A,T,C or G
W--> 150 <400> 9
W--> 151 agcggatgcc ttgtcgtcgt cgtccttgta gtctgttcgg taattaatgg aaattggsnn      60
W--> 152 snnsnnsnns nnsnnsnnsn nsnnnsnagt gacagcatat acagtgatgg tata          114
154 <210> SEQ ID NO: 10
155 <211> LENGTH: 57
156 <212> TYPE: DNA
157 <213> ORGANISM: Homo sapiens
159 <400> SEQUENCE: 10
160 agcggatgcc ttgtcgtcgt cgtccttgta gtctgttcgg taattaatgg aaattgg          57
162 <210> SEQ ID NO: 11
163 <211> LENGTH: 45
164 <212> TYPE: DNA
165 <213> ORGANISM: T7 phage and tobacco mosaic virus
167 <400> SEQUENCE: 11
168 gcgtaatacg actcactata gggacaatta ctatttaca ttaca                        45
170 <210> SEQ ID NO: 12
171 <211> LENGTH: 33
172 <212> TYPE: DNA
173 <213> ORGANISM: Artificial Sequence
175 <220> FEATURE:
176 <223> OTHER INFORMATION: Flag sequence
178 <400> SEQUENCE: 12
179 agcggatgcc ttgtcgtcgt cgtccttgta gtc                                  33
181 <210> SEQ ID NO: 13
182 <211> LENGTH: 19
183 <212> TYPE: DNA
184 <213> ORGANISM: Artificial Sequence
186 <220> FEATURE:
187 <223> OTHER INFORMATION: Splint oligonucleotide
W--> 189 <221> NAME/KEY: misc_feature
190 <222> LOCATION: (1)...(19)

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191 <223> OTHER INFORMATION: n = A,T,C or G  
 W+> 193 <221> misc\_feature  
 194 <222> LOCATION: 10  
 195 <223> OTHER INFORMATION: n = A,T,C or G  
 W+> 197 <400> 13  
 W+> 198 tttttttttt agcggatgc 19  
 200 <210> SEQ ID NO: 14  
 201 <211> LENGTH: 20  
 202 <212> TYPE: DNA  
 203 <213> ORGANISM: Artificial Sequence  
 205 <220> FEATURE:  
 206 <223> OTHER INFORMATION: Puromycin linker oligonucleotide  
 208 <400> SEQUENCE: 14  
 209 aaaaaaaaaa aaaaaaaacc 20  
 211 <210> SEQ ID NO: 15  
 212 <211> LENGTH: 30  
 213 <212> TYPE: DNA  
 214 <213> ORGANISM: Mus musculus  
 216 <400> SEQUENCE: 15  
 217 gcggcagggt ttgcttactg gggccaaggg 30  
 219 <210> SEQ ID NO: 16  
 220 <211> LENGTH: 27  
 221 <212> TYPE: DNA  
 222 <213> ORGANISM: Mus musculus  
 224 <400> SEQUENCE: 16  
 225 gggaggggtg gaggtagggtc acagtcc 27  
 227 <210> SEQ ID NO: 17  
 228 <211> LENGTH: 30  
 229 <212> TYPE: DNA  
 230 <213> ORGANISM: Mus musculus  
 232 <400> SEQUENCE: 17  
 233 tttgctagct ttaccaggag agtgggagggc 30  
 235 <210> SEQ ID NO: 18  
 236 <211> LENGTH: 33  
 237 <212> TYPE: DNA  
 238 <213> ORGANISM: Mus musculus  
 240 <400> SEQUENCE: 18  
 241 aaaaagcttg ccaaaacgac acccccatct gtc 33  
 243 <210> SEQ ID NO: 19  
 244 <211> LENGTH: 33  
 245 <212> TYPE: DNA  
 246 <213> ORGANISM: Mus musculus  
 248 <400> SEQUENCE: 19  
 249 catatgggtt ctgatattcc gagagatctg gag 33  
 251 <210> SEQ ID NO: 20  
 252 <211> LENGTH: 43  
 253 <212> TYPE: DNA  
 254 <213> ORGANISM: Mus musculus  
 256 <400> SEQUENCE: 20

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257 catatgcatc accatcacca tcacgtttct gatattccga gag 43
259 <210> SEQ ID NO: 21
260 <211> LENGTH: 30
261 <212> TYPE: DNA
262 <213> ORGANISM: Mus musculus
264 <400> SEQUENCE: 21
265 gaattcctat gttttataat tgatggaaac 30
267 <210> SEQ ID NO: 22
268 <211> LENGTH: 19
269 <212> TYPE: DNA
270 <213> ORGANISM: Homo sapiens
272 <400> SEQUENCE: 22
273 tgtaaatagt aattgtccc 19
275 <210> SEQ ID NO: 23
276 <211> LENGTH: 20
277 <212> TYPE: DNA
278 <213> ORGANISM: Homo sapiens
280 <400> SEQUENCE: 23
281 tttttttttt tttttttttt 20
283 <210> SEQ ID NO: 24
284 <211> LENGTH: 15
285 <212> TYPE: DNA
286 <213> ORGANISM: Artificial Sequence
288 <220> FEATURE:
289 <223> OTHER INFORMATION: Oligonucleotide
291 <400> SEQUENCE: 24
292 cctgtaggtg tccat 15
294 <210> SEQ ID NO: 25
295 <211> LENGTH: 16
296 <212> TYPE: DNA
297 <213> ORGANISM: Homo sapiens
299 <400> SEQUENCE: 25
300 catcgtcctt gtagtc 16
302 <210> SEQ ID NO: 26
303 <211> LENGTH: 13
304 <212> TYPE: DNA
305 <213> ORGANISM: Artificial Sequence
307 <220> FEATURE:
308 <223> OTHER INFORMATION: Oligonucleotide
310 <400> SEQUENCE: 26
311 cgtcgtaggg gta 13
313 <210> SEQ ID NO: 27
314 <211> LENGTH: 17
315 <212> TYPE: DNA
316 <213> ORGANISM: Artificial Sequence
318 <220> FEATURE:
319 <223> OTHER INFORMATION: Oligonucleotide
321 <400> SEQUENCE: 27
322 caggtcttct tcagaga 17

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RAW SEQUENCE LISTING ERROR SUMMARY  
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:3; N Pos. 74,75,77,78,80,81,83,84,86,87,89,90,92,93  
Seq#:6; N Pos. 83,84,86,87,89,90,92,93  
Seq#:9; N Pos. 59,60,62,63,65,66,68,69,71,72,74,75,77,78,80,81,83,84,86,87  
Seq#:13; N Pos. 10  
Seq#:30; N Pos. 10  
Seq#:79; Xaa Pos. 17,30,34  
Seq#:85; Xaa Pos. 28,51,82  
Seq#:96; Xaa Pos. 40  
Seq#:97; Xaa Pos. 13,21  
Seq#:107; Xaa Pos. 27  
Seq#:114; Xaa Pos. 3,18,23  
Seq#:118; Xaa Pos. 28  
Seq#:120; Xaa Pos. 28  
Seq#:125; Xaa Pos. 86  
Seq#:126; Xaa Pos. 75  
Seq#:129; Xaa Pos. 2,9,12,22  
Seq#:189; Xaa Pos. 24